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Pip Argot, Editor-in-Chief



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The Weaponized Wuhan-China Virus was No Accident

The coronavirus outbreak has brought the world to a halt.

With almost 500,000 positive cases worldwide and almost 25,000 deaths, the orchestrated COVID-19 outbreak has caused a global panic.

Rex Phineas, P.M. O.C.N. F.C.

Events are occurring now, weakening the Old World Order so that the New World Order can be established.

Rule 1: Georgia Guidestones
Maintain humanity under 500,000,000 in perpetual balance with nature.

Oi, Canadajun;

An Introduction to the Bilderberg Group

During a three-day stay in Switzerland, as part of another bid to build support for the U.S. government’s pressure campaign against Iran, American Secretary of State **Mike Pompeo** made a conspiratorial stop to a secretive **Bilderberg Group Meeting of the nefarious global elite** at Montreux—a Swiss town on the shoreline of Lake Geneva at the foot of the Alps (412 km to the Bamophetic Gotthard “Gateway to Hell”). He was not the only senior U.S. official visiting, *either*—**Jared Kushner**, President Trump’s senior adviser and son-in-law was also listed as an attendee of the powerful, above-government, occultist corporate group manipulating world events.

In 1954, the same year *that the demonic NWO Anti-Christ Beast-system* initiated their **European Organization for Nuclear Research (CERN)**—the “Stargate to Hell” in Geneva (on the Franco-Swiss border), a small group of powerful Europeans convened under leadership of **Prince Bernhard of the Netherlands**, meeting in the **Bilderberg Hotel in Oosterbeek—a village in the eastern part of Netherlands**. Its founders include exiled Polish politician **Jozef Retinger**, ex-Belgian prime minister **Paul van Zeeland** and **Paul Rijkens**, former head of consumer goods giant Unilever. This “*utopian shadow-world government*” has continued to *meet and determine our futures as “strawman” citizen-possessions* ever since... almost every year—in locations as far west as *Lake Lanier, Ga., and as far east as Istanbul*. The only year a meeting wasn’t held was in 1976, when it was canceled as Bernhard was embroiled in a scandal over accusations that he took a bribe from U.S. aircraft manufacturer **Lockheed**. [Note: **ULA (Lockheed-Boeing)** were responsible for the **Directed Free-energy Technology**—exposed by **Dr. Judy Wood**—which was responsible for the **9-11 fiasco** used to initiate a **Rothschildean bankster scheme** against the *last seven nations without one of their central banks: Libya, Iraq, Afghanistan, Sudan... Cuba, Iran and North Korea* are yet incompliant.]

The known list of past attendees is vast. A number of them have gone on to lead nations shortly afterward: The disgusting “*pinko occultist*” **Angela Merkel** attended in 2005, only months before becoming the German chancellor, while *drug-dealing “Rhodes scholar” libertine Bill Clinton* attended in 1991 when he was still governor of Arkansas. It is relatively unusual for a serving U.S. secretary of state like Pompeo to attend, though not unprecedented: **Condoleezza Rice** was listed as a participant in 2008, when the event was held in Chantilly, Va., while she was still head of the State Department. Nor are Pompeo and Kushner the first Trump administration staffers to attend: In 2017, when the event was held again in Chantilly, administration officials including Commerce Secretary **Wilbur Ross** and national security adviser **H.R. McMaster** attended.

About 130 *ruthless pan-global fascists* and *totalitarian Cultural Marxists* attend the *shadowy conference of international power brokers* every year, with two-thirds generally from Europe and one-third from North America. A third are in government and politics, and the rest are from other fields—corporate, military and banking—its members plotting **New World Order**, hell-bent on *world domination* and reinforcing their interests by choosing *presidents, politicians* and *captains of industry*—as *the unelected European Union parliamentarians*, controlling public opinion through *puppet media magnates* which propagate false news claiming the Bilderberg Group is nothing more sinister than “an occasional supper club.”

The 2018 list of invitees included Canadian Minister of International Trade **François-Philippe Champagne** and Quebec’s deputy premier **Geneviève Guilbault**. Former Bank of Canada Governor **Mark Carney**, currently the governor of the Bank of England, was *also* listed as a participant, along with billionaire **Peter Thiel**, Secretary General of NATO **Jens Stoltenberg**, the **King of the Netherlands**, a smattering of *prominent newspaper editors, journalists and professors*, and *Prime Ministers from Belgium, Estonia and the Netherlands*.

The 2019 attendee list included many core participants, such as former secretary of state and national security adviser **Henry Kissinger**. The war criminal Kissinger is a regular, while **Helmut Kohl**, **Bill Gates**, **Christine Lagarde** and **Jose Manuel Barroso** have all attended among the *billionaires and executives from leading banks, corporations, defense industry bigwigs* and their allies on **Wall Street** and in **Silicon Valley**; operating behind the scenes and *manufacturing “hive-mind” public opinion* via news media, academia, and the sports and entertainment industry.

Participants at Bilderberg Meetings analyze and promote major international issues and current affairs but are barred from revealing who said what during discussions under **Chatham House Rules**—as an aid to free discussion of sensitive issues. **Chatham House**—or, the Rothschildean **Royal Institute of International Affairs**—an independent London-based “*Crown City*” *policy-making institute* affiliated with Rockefeller’s **Council on Foreign Relations** and the **Trilateral Commission**. Speakers openly discuss their views in private while allowing the generalized topics and nature of the debates to be made public... but the group does not release any notes or summaries of discussions: Notwithstanding, as a gesture towards transparency, the *sinister cabal of the rich and powerful* with something to hide publishes its guest list the day before its annual get together—between 120-and-150 of their most affluent and propertied devilish puppets are invited by its steering committee—a generalized list of what subjects they intend to discuss is also released for public scrutiny, vaguely consisting of broad issues and macroeconomic concerns, such as the threat of *terrorism and cyber-security*.

Journalists are strictly forbidden to report on the event, though some *may* attend as participants. More-so, such partakers may freely make use of any information they are programmed with, but neither the identities nor affiliation of the speakers, nor what is said or done of any other participant, may be revealed.

A Brief History of SARS-CoV-2

Official names have been announced for the virus responsible for COVID-19, previously known as “**novel coronavirus disease-2019**”—the virus is referred to as “**severe acute respiratory syndrome coronavirus 2**” (SARS-CoV-2).

Viruses, and the diseases they cause, often have different names. For example, **HIV** is the virus that causes **AIDS**. People often know the name of a disease, but not the name of the virus that causes it. There are different processes, and purposes, for naming viruses and diseases. Viruses are named based on their *genetic structure* to facilitate the development of diagnostic tests, vaccines and medicines. Virologists and the wider scientific community do this work, so viruses are named by the **International Committee on Taxonomy of Viruses (ICTV)**.

Diseases are named to enable discussion on disease prevention, spread, transmissibility, severity and treatment. Human disease preparedness and response is the responsibility of the **World Health Organization (WHO)**, so diseases are officially named by WHO in the **International Classification of Diseases (ICD)**. On 11 February 2020, ICTV announced “**severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2)**” as the name of the new virus—and, following guidelines previously developed with the **World Organization for Animal Health (OIE)** and the **Food and Agriculture Organization of the United Nations (FAO)**, WHO announced “**COVID-19**” as the name of this new disease. This name was chosen because the virus is genetically related to the coronavirus responsible for the **SARS outbreak of 2003**. While related, the two viruses are different.

Middle East Respiratory Syndrome (MERS)

Middle East Respiratory Syndrome (MERS) also known as “*camel flu*,” is a viral respiratory disease caused by a **lineage C Betacoronavirus** that emerged in the human population in 2012. Since 2012, the **World Health Organization (WHO)** has reported 2266 confirmed human cases, including 804 deaths. **Middle East Respiratory Syndrome Coronavirus (MERS-CoV)** is a *zoonotic virus* that has repeatedly entered the human population *via* direct or indirect contact with infected dromedary camels in the Arabian Peninsula. First reported in **Saudi Arabia** in 2012, it has since spread to the U.S.: Globally, MERS is active in five other countries: **Malaysia, Oman, United Arab Emirates, United Kingdom of Great Britain and Northern Ireland**, as well as the **Republic of Korea**. The most likely threat to Canadians remains a risk of importation to Canada from immigrants infected while in the Middle East... but, as of yet, no cases of MERS have been reported in Canada.

WHO and colleagues from the **University of Oxford, Imperial College London** and **Institut Pasteur** have estimated that, since 2016, 1465 cases of Middle East Respiratory syndrome coronavirus (MERS-CoV) and between 300 and 500 deaths may have been averted due to accelerated global efforts to detect infections early and reduce transmission. The risk of exported cases to areas outside of the Middle East due to *immigration* remains significant. At the end of November 2019, a total of 2494 laboratory-confirmed cases of Middle East respiratory syndrome (MERS), including 858 associated deaths (case–fatality rate: 34.4%) were reported globally; the majority of these cases were reported from Saudi Arabia (2102 cases, including 780 related deaths with a case–fatality rate of 37.1%).

Source of Zoonotic Transmission

Research to date suggests that MERS-CoV emerged from bats (Sharif-Yakan & Kanj, 2014). Genomic sequencing and phylogenic analysis from bats in Saudi Arabia and Africa support this theory, with similarities identified between bat coronavirus sequences and MERS-CoV (Ithe et al., 2013; Memish et al., 2013; Wang et al., 2014). Multiple studies on the emergence and transmission of MERS-CoV point to the role of dromedary camels as a reservoir and primary source of zoonotic infection due to (1) the high prevalence of MERS-CoV antibodies in camels across the Arabian Peninsula, North and Eastern Africa, (2) the detection of MERS-CoV by RT-PCR in respiratory secretions, feces, and milk of dromedary camels and (3) similarities in the genomic sequence of viruses isolated from infected individuals and camels (Omrani, Al-Tawfiq, & Memish, 2015). There is evidence to suggest that MERS-CoV has been circulating in camels for more than 20 years (since at least 1992) (Alagaili et al., 2014; Corman et al., 2014). Challenge studies conducted in camels suggest that MERS is a mild respiratory infection resulting in little to no signs of illness in camels (Adney et al., 2014). Based on seroepidemiological studies carried out in other livestock, alpacas are the only other livestock in which MERS-CoV specific antibodies have been identified; however, no human cases with exposure to alpacas have been reported to date (Reusken et al., 2016).

Although the route of camel to human transmission is not fully understood, zoonotic infections of MERS-CoV likely occur through direct or indirect contact with infected camels or camel-related products (e.g. raw milk) (Gossner et al., 2016). The detection of MERS-CoV RNA in camel milk suggests a potential food-borne transmission of the virus by consuming raw milk (Reusken et al., 2014). Epidemiological studies have demonstrated a greater risk of MERS-CoV infection among individuals exposed to camels when compared to the general population (Muller M.A. et al., 2015; World Health Organization [WHO], 2015b). To date, zoonotic transmission has only been reported from countries in the Arabian Peninsula where the virus is highly prevalent and widespread within the camel population. Further studies need to be conducted in order to fully understand the role human-animal interaction play in the transmission of this disease.

Source of Secondary Transmission

Secondary transmission of MERS-CoV has largely been associated with nosocomial outbreaks; however, family clusters have also been reported (Abroug et al., 2014; Memish, Zumla, Al-Hakeem, Al-Rabeah, & Stephens, 2013). Nosocomial outbreaks of MERS have played a significant role in increasing the magnitude of reported cases. To date, large scale hospital outbreaks have been reported in Saudi Arabia and South Korea. Lack of IPC (e.g., overcrowding, delay in isolation of cases, lack of knowledge of MERS case definitions amongst healthcare workers) along with intra- and inter-hospital transmission, super spreaders, and asymptomatic cases are a few of the factors that have contributed to these hospital outbreaks (Al-Tawfiq & Perl, 2015). Strict adherence to IPC has been shown to result in the rapid identification of cases and a significant reduction in the number of secondary cases in nosocomial outbreaks, and allows for early identification of new cases (El Bushra et al., 2017).

An outbreak in South Korea, which began in May 2015, was the largest nosocomial outbreak of MERS outside the Middle East. This outbreak started from a super spreader who traveled to four countries in the Middle East and who appeared to have transmitted infection to close relatives, patients sharing the same room and health care workers providing care upon return to South Korea (Kim et al., 2017). These exposures happened before MERS-CoV was suspected or diagnosed. The outbreak in the Republic of Korea was declared over on December 23, 2015 (Normile, 2015). The current evidence from contact tracing studies suggests that transmission did not extend beyond close contacts into the community (Memish, Zumla, & Assiri, 2013)

Vaccines against MERS-CoV

No vaccine or effective antiviral treatment is currently available for MERS-CoV. However, research is underway and many candidate vaccines targeted for humans are in development. These candidates use a wide range of platforms including whole virus vaccine (live attenuated and inactivated), vectored-virus vaccines, DNA vaccine, and

protein-based vaccines (Okba, Raj, & Haagmans, 2017). Thus far, only one candidate vaccine has advanced to clinical development: a DNA-only vaccine that has been shown to generate protective MERS-CoV antibodies in mice, camels and monkeys advanced to Phase 1 human clinical trials in February 2016 (Modjarrad, 2016).This candidate vaccine has also been tested in dromedary camels. Additionally, a modified vaccinia Ankara based vaccine that reduced virus shedding and generated protective antibodies and mucosal immunity is currently scheduled for a Phase 1 safety trial (Alharbi, 2017).

Virus Characteristics

MERS-CoV is an enveloped and single stranded RNA virus belonging to the Coronaviridae family (Fehr et al., 2017). Though caused by the same genera of coronavirus as severe acute respiratory syndrome (SARS) (Betacoronavirus genera), MERS-CoV belongs to lineage C whereas SARS belongs to lineage B (Arabi Y.M. et al., 2017). Entry of MERS-CoV into host cells is mediated by its spike (S) protein binding to the host cell receptor, dipeptidyl peptidase 4 (DPP4) (Fehr et al., 2017). In the case of SARS, extensive mutations in the virus resulted in its ability to adapt to the human cell receptor angiotensin-converting enzyme 2 [ACE2] which allowed for increased human-to-human transmission (Chinese SARS Molecular Epidemiology Consortium, 2004). Though sequenced viral isolates from camels and humans indicate that MERS-CoV has undergone recombination and genetic changes since its introduction to humans in 2012 (Sabir et al., 2016), none of these changes have resulted in virus mutation allowing for more efficient adaptation to human cell receptor, DPP4. In fact, analysis of viral isolates from the South Korea outbreak with a mutation in the receptor binding domain showed reduced affinity to human DPP4 (Kim et al., 2016). These findings further convolute the role of the S protein and DPP4 in the human adaptation of this virus given the magnitude of the South Korea outbreak.

Republican senator Tom Cotton of Arkansas

Republican senator **Tom Cotton** of Arkansas, a foreign policy hardliner, suggested “the coronavirus was manufactured in a ***Chinese bioweapons level-four Cellular Level Biosafety super-laboratory researching human infectious diseases.***” In short, Cotton is suggesting that either “the virus is the result of an experiment gone badly” or “a purposeful release of a bio-weapon.” Since January, the first-term senator has warned that China was covering up the lethality of the disease and doing little to stop what ha become ***a global pandemic.*** In December 2016, **Bill Gates** warned **Donald Trump** before he took office of the dangers of a pandemic—and urged him to prioritize the America’s preparedness efforts. Cotton, other leading Trump allies and the president himself were mocked, despite not making a serious accusation it was intentional, referring to Covid-19 as the **Wuhan or Chinese virus**—and denying that it is racist or discriminatory to do so.

“Everyone with whom I’ve spoken, or whom I’ve read, thinks that Covid-19 has come from a natural source, as did the SARS virus, as did the MERS virus. Both of those were also coronaviruses in animal populations that jumped to the human species in the natural environment.”

Dr. William Schaffner, infectious disease specialist, Vanderbilt University Medical Center

But burden of proof is on Chinese Ambassador to the United States **Cui Tiankai** and his fellow communists.

“China’s criminal negligence allowed what could have been a health outbreak in Wuhan to become a devastating global pandemic. Their malignant and deliberate action sent the virus around the world by allowing international travel to continue in December and January, and represents just how little regard for human life they have.”

Republican senator **Tom Cotton**

In April, **Yuan Zhiming**, secretary of the **Chinese Communist Party committee at Wuhan Institute of Virology** and a top researcher at the lab—linked to China’s covert biological weapons program, which studies some of the world’s deadliest pathogens. He denied his laboratory was a possible origin for the pandemic—while ***the Chinese government denied the possibility of a lab leak,*** its actions, *though*, tell a different story: The Chinese military posted its top epidemiologist to the Institute of Virology in January. In February **Chairman Xi Jinping** urged ***swift implementation of new biosafety rules to govern pathogens in laboratory settings.*** Academic papers about the virus’ origins are now subject to prior restraint by the government.

“Now, we don’t have evidence that this disease originated there, but because of China’s duplicity and dishonesty from the beginning, we need to at least ask the question to see what the evidence says. And China right now is not giving any evidence on that question at all.”

Republican senator **Tom Cotton**

The early scientific consensus was that the ***never-before-seen coronavirus*** had most likely made a jump from ***bats*** to another form of wildlife, perhaps the type of exotic animal, like a ***pangolin***, sold at **Wuhan’s Huanan Seafood Wholesale Market**—an unregulated meat market full of wild animals, where it then achieved a ***natural zoonotic transmission to humans.*** But scientists don’t rule out that an accident at a research laboratory in Wuhan might have spread ***a deadly bat virus*** that had been collected for scientific study. Less than 300 yards from the seafood market is the Wuhan branch of the **Chinese Center for Disease Control and Prevention.**

“It will not be possible to assign a numerical probability to the lab-accident scenario without a forensic investigation. A forensic investigation would require access to lab facilities, samples, records, and personnel, and would include environmental sampling of facilities and serological sampling of personnel. A forensic investigation—if one occurs—would face the same challenges as the 2001–2008 investigation of the 2001 anthrax mailings in the U.S., plus additional challenges arising from more time to alter evidence and from dealing with a government not known for transparency and introspection.”

Richard Ebright, biosecurity expert, Waksman Institute of Microbiology, Rutgers University

Bottom line: Nobody knows for sure. The type of investigation necessary to establish more certainty would require cooperation not likely to be found in China. Thanks to the ***Chinese cover-up,*** we may never have direct, conclusive evidence of ***the source of the coronavirus:*** In early-January Red Chinese enforcers threatened doctors who warned their colleagues about the virus; among them was **Li Wenliang**, who died of Covid-19 in February. Laboratories working to sequence ***the virus’ genetic code*** were ordered to destroy their samples. The laboratory that first published the ***virus’ genome*** was shut down, Hong Kong’s **South China Morning Post** reported in February.

Cotton’s remarks were not the first time it was suggested that the virus may have originated in the Wuhan lab. The British publication the **Daily Mail** was one of the first to suggest a connection between the coronavirus and the laboratory in Wuhan. Later, the **Washington Times** ran a story under the headline, “Coronavirus may have originated in lab linked to China’s biowarfare program.”

“In around 2020 a severe pneumonia-like illness will spread throughout the globe, attacking the lungs and the bronchial tubes and resisting all known treatments.”

“End of Days: Predictions and prophecies about the end of the world” by Sylvia Browne and Lindsay Harrison (2008)

The disease isn’t deadly enough to kill off the Chinese customer base, nor the customers who buy products from **Silicon Valley** or from Europe. But, with the aid of a complaint media (*q.v.*) it won’t be portrayed as anything sinister, even if it suddenly slams the brakes on the worldwide economy. The American expert on biological threats **Francis Boyle** confirmed the suspicions of many, pointing to the BIs-4 laboratory in Wuhan, where the epidemic broke out. What he doesn’t mention are the 25 military centers for testing

epidemic viruses open all over the world by the **Pentagon Defense Threat Reduction Agency (DTRA)**—as noted in 2018 by Bulgarian journalist **Dilyana Gaytandzhieva.**

In her investigation, the reporter *also* described tests on ***bio-genetic weapons capable of affecting a specific ethnic group*** and considered ***“potentially unstoppable”*** by a study published in August 2019 by the **Center for the Study of Existential Risk (CSER)** of the University of Cambridge. A suspicious CoronaVirus research was also launched in 2015 by the British **Pirbright Institute** funded by the **Melinda and Bill Gates Foundation.**

Theories about a connection between the Pirbright Institute and Bill Gates and the novel coronavirus have been circulating online for months. While the **Pirbright Institute does hold a patent for a coronavirus** *that primarily affects chickens*, it claims not to hold any patents on parts of or the complete ***genome of the SARS-CoV-2 betacoronavirus*** causing the current COVID-19 pandemic.

Avian (bird) Influenza (flu) Type A

The Melinda and Bill Gates Foundation, big contributors to the Bilderberg Group and proponents of ***deadly vaccines,*** did fund a research center in England called the **Pirbright Institute**—a publicly funded private company which specializes in “the study of viruses that affect farm animals,” and “viruses which transfer from animals to people,” but ***they do not own a patent on the COVID-19 coronavirus.***

Remember, it was just over a dozen years ago that ***a bird flu virus*** known as **H5N1** was charting a destructive course through Asia, North Africa, and the Middle East, ravaging poultry in apocalyptic numbers and killing 6-in-10 humans known to have contracted it—scientists and government officials feared that the virus could ignite a human pandemic reminiscent of the catastrophic **1918 Spanish flu.**

The virus continued to kill chickens and to *occasionally* infect and sometimes kill people. Currently, the avian influenza H5N1 virus continues to circulate in poultry in some countries, especially in Asia and northeast Africa. This virus continues to cause ***sporadic human infections*** with some instances of limited human-to-human transmission among very close contacts. There has been no sustained human-to-human or community-level transmission identified thus far. But as the years passed, the number of ***human H5N1 cases*** subsided. There has not been a single H5N1 human infection detected since February 2017. The virus’ nearly-as-scary cousin, **H7N9**, emerged in China in 2013 and sickened more than 1500 people in China over five years, killing roughly 40% of them. But after an extraordinary surge of 766 cases in early 2017, there were a mere three infections recorded in 2018; in 2019 there were none. While bird flu may no longer be in the headlines, it very much still remains a concern for influenza experts

“The H5N1 virus has not gone away. It’s just changed into different versions of itself.” Malik Peiris, professor of virology and influenza expert (University of Hong Kong)

During the post-peak period, pandemic disease levels in most countries with adequate surveillance will have dropped below peak observed levels. The post-peak period signifies that pandemic activity appears to be decreasing; however, it is uncertain if additional waves will occur and countries will need to be prepared for a second wave. Previous pandemics have been characterized by waves of activity spread over months: Once the level of disease activity drops, a critical communications task will be to balance this information with the possibility of another wave. Pandemic waves can be separated by months and an immediate “at-ease” signal may be premature.

In the post-pandemic period, influenza disease activity will have returned to levels normally seen for seasonal influenza. It is expected that the pandemic virus will behave as a seasonal influenza A virus. At this stage, it is important to maintain surveillance and update pandemic preparedness and response plans accordingly. An intensive phase of recovery and evaluation may be required.

The Pirbright Institute carries out research on **infectious bronchitis virus (IBV)**, a coronavirus that infects poultry and **porcine deltacoronavirus** that infects pigs. Pirbright, *though*, claims “it does not currently work with human coronaviruses.”

The **Center for Disease Control and Prevention (CDC)** claims that there is already a vaccine for a certain strain of **Ebola**, but does not say if a vaccine is available for **Zika** and/or COVID-19. Yet, even if there were a vaccine it would contain the virus. After getting a vaccine, the imitation infection can cause symptoms, such as fever, and should be expected as the body builds immunity; so... anyone getting vaccinated—*causing the immune system to produce T-lymphocytes and antibodies*, might become infected with the virus.

The D614G Mutation

A mutation in the protein that allows SARS-CoV-2 to enter cells will make it easier for the virus to spread—but the but the lying public health officials are saying it might not make a difference at all. The mutation is not new. It appears in low levels in samples taken from COVID-19 patients as far back as February. But this variation of the virus (nicknamed the “G” variation) seems to show up in more and more of the virus samples taken from people infected recently compared to early in the pandemic.

The Coronavirus was Predicted by the Georgia Guidestones

The **World Health Organization** lists more than 30 ***potential vaccines*** currently in development for treating COVID-19. Much of the research being conducted by pharmaceutical companies around the world, who will seek to patent a vaccine upon its completion and put a price-tag on it for folks like us—an out-of-pocket expense for the working poor and paid for by tax dollars for those on social assistance.

In 2017, **Jeremy Konyndyk**, former director of the **United States Agency for International Development (USAID)**’s Office of **US Foreign Disaster Assistance** under the Obama administration; and, in 2018, the intelligence community’s **Worldwide Threat Assessment** warned that a “novel strain of a virulent microbe which was easily transmissible between humans continues to be a major threat.” The 2019 threat assessment from January stated: “We assess [] the world will remain vulnerable to the next flu pandemic or large-scale outbreak of a contagious disease that could lead to massive rates of death and disability, severely affect the world economy, strain international resources.”

In 2009, after years of studying infectious diseases at the CDC and USAID, **Dennis Carroll** saw that emerging infectious diseases, far and wide, have mostly come from wildlife... and humans were proximal. The first thing to understand is that whatever future threats we’re going to face already exist. A large pool of viruses is even now circulating in the wild—and we don’t become familiar with them until we see a spillover event and people begin getting ill.



The “cornea” is a clear window at the front of the eye.

Doomsday warnings are spreading, blending coronavirus fears with everything from political paranoia to references from the Bible. Predictions of apocalyptic events that would result in the extinction of humanity, a collapse of civilization, or the destruction of the planet have been made since the origin of history. Religious wars, global terrorism, pandemics, and genocide have all helped to usher in the Anxiety Age.

O Canada,
God Bless and Be Well.

o/